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Evidence Appendix

A copy of the article by Prasher et. al entitled "Primary Structure of the Aequorea Victoria Green Fluorescent Protein", Gene 111: 229-233 (1992) is appended hereto. This article was first cited by the Examiner in an Office Action dated March 14, 2003.

A copy of U.S. Patent No. 5,246,631 to Halbritter is appended hereto. This patent was first cited by the Examiner in an Office Action dated March 14, 2003.

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Primary structure of the *Aequorea victoria* green-fluorescent protein

(Bioluminescence; Cnidaria; aequorin; energy transfer; chromophore; cloning)

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SUMMARY

Many cnidarians utilize green-fluorescent proteins (GFPs) as energy-transfer acceptors in bioluminescence. GFPs fluoresce in vivo upon receiving energy from either a luciferase-oxyluciferin excited-state complex or a Ca^{2+} -activated photoprotein. These highly fluorescent proteins are unique due to the chemical nature of their chromophore, which is comprised of modified amino acid (aa) residues within the polypeptide. This report describes the cloning and sequencing of both cDNA and genomic clones of GFP from the cnidarian, *Aequorea victoria*. The *gfp10* cDNA encodes a 238-aa-residue polypeptide with a calculated M_r of 26888. Comparison of *A. victoria* GFP genomic clones shows three different restriction enzyme patterns which suggests that at least three different genes are present in the *A. victoria* population at Friday Harbor, Washington. The *gfp* gene encoded by the λ GFP2 genomic clone is comprised of at least three exons spread over 2.6 kb. The nucleotide sequences of the cDNA and the gene will aid in the elucidation of structure-function relationships in this unique class of proteins.

INTRODUCTION

Luminescence is common in a variety of marine invertebrates. Many cnidarians and probably all ctenophores emit light when mechanically disturbed. Proteins responsible for bioluminescence from several species of these two

phyla have been characterized. Light from luminescent cnidaria is primarily green whereas light emitted from ctenophores is blue. The green light of cnidaria is due to the presence of a class of proteins called green-fluorescent proteins (GFPs). They are highly fluorescent and are activated in vivo by an energy transfer process via a luciferase or a Ca^{2+} -activated photoprotein, both of which produce energy during the oxidation of coelenterate-type luciferin. In the cnidarian *Aequorea*, the photoprotein aequorin excites the GFP by an unknown mechanism to release green light. Previous studies suggesting that *Aequorea* GFP is stimulated via a radiationless mechanism (Morise et al., 1974) have been questioned (Ward, 1979). The GFP from *Renilla*, another cnidarian, on the other hand, clearly receives energy from the *Renilla* luciferase-oxyluciferin excited state.

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Abbreviations: *A.*, *Aequorea*; aa, amino acid(s); bp, base pair(s); GFP, green-fluorescent protein; *gfp*, DNA or RNA encoding GFP; kb, kilobase(s) or 1000 bp; nt, nucleotide(s); oligo, oligodeoxyribonucleotide; ORF, open reading frame(s).

complex by a radiationless energy transfer mechanism (Ward and Cormier, 1976).

The GFPs most thoroughly studied have been isolated from *Aequorea* and *Renilla* (Ward, 1979). The *Aequorea* GFP has been reported to be a 30-kDa monomer (Pren-dergast and Mann, 1978) whereas the *Renilla* GFP is a 54-kDa homodimer (Ward and Cormier, 1979). The two proteins have different absorption spectra but identical emission spectra ($\lambda_{\text{max}} = 509 \text{ nm}$). Upon denaturation the two GFPs have the same absorption spectra. Ward et al. (1980) have predicted that both *Aequorea* and *Renilla* GFPs contain chromophores having the same structure but that the different absorption spectra are explained by different apoprotein environments.

Biochemical properties of the *Aequorea* GFP show it to have unique structural properties. The fluorescent chromophore is stable to a variety of harsh conditions including heat, extreme pH, and chemical denaturants. Fluorescence is lost, for example, to base or acid treatment or addition of guanidine hydrochloride, but upon neutralization of the pH or removal of the denaturant, fluorescence returns with an identical emission spectrum (Bokman and Ward, 1981; Ward and Bokman, 1982). The chromophore structure is very different from those of the phycobiliproteins which are also highly fluorescent. The chromophore in the GFPs is covalently bound and is formed by modification of certain aa residues within the polypeptide. The chemical structure of the *Aequorea* GFP chromophore (Fig. 1), first characterized by Shimomura (1979), has been thoroughly re-examined (Ward et al., 1989; W.W.W., unpublished) and is shown here (Fig. 1) in its revised form. In this study, the *Aequorea* GFP gene and its cDNA have been isolated and characterized in pursuit of elucidating the mechanism of energy transfer between aequorin and GFP as well as addressing evolutionary relationships in coelenterate bioluminescence.

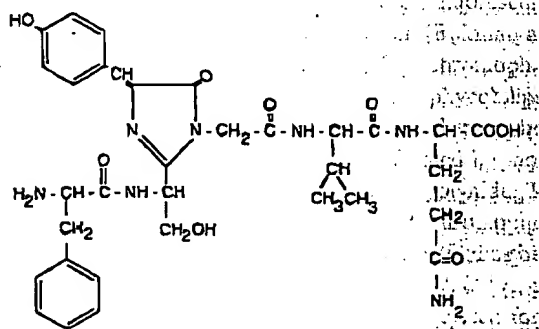


Fig. 1. The chemical structure of the chromophore in *Aequorea* GFP (W.W.W., unpublished). The cyclized chromophore is formed from the trimer Ser-dehydroTyr-Gly within the polypeptide by an unknown mechanism.

EXPERIMENTAL AND DISCUSSION

(a) Construction of cDNA libraries

An *A. victoria* cDNA library, constructed in pBR322 (Prasher et al., 1985), was screened for the presence of a *gfp* cDNA using two oligo mixtures whose sequences were based on the aa sequences derived from GFP-derived CNBr fragments. The oligos contained the following nt sequences: A: 5'-AA₆AA₆TC₆TG₆TG₆TTTCAT (20-mer with 32 redundancies), B: 5'-TT₆TA₆TT₆TA₆TCCAT (17-mer with 16 redundancies). The hybridization of the ³²P-labeled mixtures A and B to replicate filters containing this library were performed according to the method of Wood et al. (1985) utilizing tetramethylammonium chloride during the washing steps. The temperatures used during the washing steps for mixtures A and B were 55°C and 50°C, respectively.

A single *gfp* cDNA was isolated from the library by this method. This clone, pGFP1, contained a *Pst*I insert of 511 bp having an ORF encoding 168 aa. The deduced translation of the nt sequence indicated the *gfp1* cDNA lacked both the 5'- and 3'-sequences of the coding region. However, the sequence FSYGVQ within the deduced translation permitted the chromophore structure to be deciphered (W.W.W., unpublished). Upon rescreening the library with *gfp1* cDNA, no additional cDNAs were found.

A second *A. victoria* cDNA library was constructed (Gubler and Hoffman, 1983) in λ gt10 (Huynh et al., 1985). The *Pst*I insert from *gfp1* cDNA was used as a hybridization probe against the entire λ gt10 library of 1.4×10^6 recombinant phage. No *gfp*-related recombinants were identified upon screening the primary library. The phage remaining on the plates were extracted from the top agar and used as an amplified library (Maniatis et al., 1982). Upon screening this preparation of the library, four recombinants hybridized to the *gfp1* cDNA following their purification. The four cDNA clones were designated λ GFP10, 11, 12, and 13. All four recombinants were shown to contain an insert of 1 kb upon digestion with *Eco*RI.

(b) Characterization of the *gfp10* cDNA

The entire *Eco*RI insert of λ GFP10 was sequenced (Fig. 2). Limited nt sequences obtained from λ GFP11 and 12 were identical with that from λ GFP10 suggesting that they were siblings and, hence, were not sequenced further. Even though the entire coding region appears to be present (see below), three features of the cDNA insert of λ GFP10 suggest it is not quite full-length. First, the cDNA is 965 nt where the *gfp* mRNA is 1.05 kb in length as determined by Northern analysis (Fig. 3). Second, the 5'-untranslated region is very short. Third, no poly(A) track is observed in the *gfp10* cDNA sequence (Fig. 2) despite the presence of the *gfp* mRNA in only the poly(A)⁺ RNA fraction of *A. vic-*

ATACACCAAA	TAAGACATAA	CAAAAG	ATC	ACT	AAA	GGA	GAA	CAA	CTT	TTC	ACT	GGA	GTT	GTC	CCA	ATT	CTT	GTT	CAA	TGA	CAT	GCT	85					
N	S	K	G	R	L	P	T	G	V	V	P	I	L	V	E	L	D	C					20					
CAT	GTT	AAT	GGG	CAC	AAA	TTT	TCT	CTC	ACT	GCA	GAG	GCT	CAA	GCT	CCA	ACA	TTC	GCA	AAA	CTT	ACC	CTT	AAA	TTT	ATT	166		
D	L	V	N	G	R	P	S	V	S	C	E	G	E	G	D	A	T	F	C	K	L	T	L	K	F	Z	47	
TCC	ACT	ACT	GCA	AAA	CTA	CCT	GTT	CCA	TGG	CCA	ACA	CTT	CTC	ACT	ACT	TTC	TCT	TAT	GCT	GTT	CAA	TCC	TTT	TCA	ACA	TAC	247	
C	T	T	G	K	L	P	V	P	M	P	T	L	V	T	T	P	S	Y	G	V	Q	C	F	S	R	Y	74	
CCA	CAT	CAT	ATC	AAA	CAG	CAT	CAC	TTT	TTC	AAG	AGT	GCC	ATC	CCC	GAA	GCT	TAT	GTA	CAG	GAA	ACA	ACT	ATA	TTT	TTC	AAA	328	
F	D	B	H	K	Q	E	D	F	F	K	S	A	H	P	E	G	Y	V	O	E	R	T	I	F	F	K	101	
GAT	CAC	GGG	AAC	TAC	AAG	ACA	CCT	GCT	CAA	GTC	AAG	TTT	GAA	GCT	CAT	ACC	GTT	GTT	AAT	AGA	ATC	CAC	TGA	AAA	GCT	ATT	409	
D	D	G	N	Y	K	T	R	A	E	V	K	P	E	C	D	T	L	V	N	R	I	E	L	K	G	I	128	
CAT	TTT	AAA	CAA	CAT	GCA	AAC	ATT	CTT	GGA	CAC	AAA	TTG	GAA	TAC	AAC	TAT	RAC	TCA	CAC	AAT	GTA	TAC	ATC	ATC	GCA	CAC	490	
D	F	K	E	D	G	N	I	L	G	N	K	L	E	Y	N	T	N	S	H	N	V	Y	I	M	A	D	155	
AAA	CAA	AAC	AAT	GGA	ATC	AAA	CTT	AAC	TTC	AAA	ATT	ACA	CAC	AAC	ATT	CAA	CAT	CCA	AGC	GTT	CAA	CTA	GCA	CAC	CAT	TAT	571	
K	Q	K	N	G	I	K	V	N	P	R	I	R	H	N	I	E	D	G	S	V	Q	L	R	D	H	Y	182	
CNA	CAA	AAT	ACT	CCA	ATT	GCC	CAT	GGC	GCT	CTC	CTT	TGA	CCA	CAC	AAC	CAT	TAC	CTC	TCC	ACA	CAA	TCT	GCC	CTT	TGC	AAA	652	
Q	Q	N	T	P	I	G	D	G	P	V	L	L	P	D	N	H	I	L	S	T	Q	E	A	L	S	K	209	
GAT	CCC	AAC	GAA	AAG	ACA	CAC	CAC	ATG	GTC	CTT	CIT	CAG	TTT	GTA	ACA	GCT	GCT	GCC	ATT	ACA	CAT	GCC	ATG	GAT	GAA	CTA	733	
D	F	N	E	K	R	D	E	H	V	L	L	E	P	V	T	A	A	G	I	T	B	Q	H	D	E	L	236	
TAC	AAA	TAA	ATGTCACAC	TTCCAAATTA	CACATAAGTC	TCGACAAAT	TACTAAATC	TCAGGCTTC	TGCTTAAT	CAGGCTCA	TATATTTAT	832															238	
Y	K																											
ATATTATAG	ATTCATATAA	ATTGATGAA	TAAATTATTC	ATGTTATTA	TAGAGCTTAT	TTTCTATTA	AACAGCTTAC	TTGAGCTGA	TTCTTAATC	932																		
TATATTAAAT	ACATTTGAT	TTCACTTCT	CAAA	962																								

Fig. 2.

Fig. 2. Nucleotide sequence of the *gfp10* cDNA and the deduced aa sequence. Below the first nt of each codon is the single-letter designation for the aa. The horizontal lines underline those aa sequenced directly from native GFP. The downward arrows indicate the positions of introns when compared to the nt sequence of the *gfp2* gene. Arrowhead: start codon; period: stop codon. DNA fragments from both cDNA and genomic clones were subcloned into M13mp18 and M13mp19 (Yanisch-Perron et al., 1985), and unidirectional deletions were prepared using the method of Dale et al. (1985). Sequencing was performed using either the Klenow fragment or an altered T7 DNA polymerase (Sequenase Ver 2.0, United States Biochemical Corp.) in the dideoxy chain termination method (Sanger et al., 1977). Both DNA strands of the sequences described in this report here have been sequenced. The GenBank accession No. for the *gfp10* sequence is M62653.

Fig. 3. Northern analysis of the *A. victoria gfp* mRNA. The poly(A)⁺ mRNA (lane 1) was denatured using glyoxal prior to electrophoresis, as described by Thomas (1983). Electrophoresis was performed for 3 h in a 1% agarose gel (pretreated with 10 mM sodium iodoacetate) equilibrated in 10 mM sodium phosphate pH 7.0 buffer. Overnight transfer of the nucleic acids to nitrocellulose was facilitated with 20 × SSC. Hybridization of ³²P-labeled *gfp1* cDNA to the membrane-bound nucleic acids was at 42°C for 28 h in 5 × SSC/5 × Denhardt's/20 mM Na₂ phosphate pH 6.8/100 µg per ml of denatured herring sperm DNA/10% polyethyleneglycol/50% formamide. HindIII-digested λ DNA, ³²P-labeled, and treated in parallel with the RNA, was used as molecular weight standards (lane 2).

tor RNA (data not shown). A typical polyadenylation signal is located at nt 861–865 (Fig. 2).

The nt sequence of the *gfp10* cDNA contains an ORF encoding a 238-aa protein having a calculated *M_r* of 26888. This compares favorably with 30 kDa for native GFP as determined by denaturing electrophoresis (Prendergast and Mann, 1978). The deduced translation contains aa sequences of numerous peptides isolated from native GFP (underlined in Fig. 2). When compared to the *gfp10* cDNA sequence (Fig. 2), the *gfp1* cDNA was determined to encode aa residues 28–195. Oligo mixture A is complementary to the codons encoding aa 78–84 and mixture B is complementary to the codons encoding aa 141–146 (Fig. 2). The trimer Ser-Tyr-Gly, modified in the native protein to form the chromophore (W.W.W., unpublished), is located at aa 65–67. The chromophore consists of an imidazolone ring formed by the residues Ser-dehydroTyr-Gly within the

polypeptide (Fig. 1). Located 8 aa upstream of this chromopeptide is GFP's only Trp. The inability to detect the fluorescence from this Trp makes it unusual (W.W.W., unpublished). Perhaps energy-transfer occurs between it and the chromophore in the native protein preventing the Trp fluorescence (320–350 nm). The Trp is flanked by several Pro residues (Pro-Val-Pro-Trp-Pro). The significance of this pentapeptide is not understood but a search of the protein databases (PIR ver 25; Swiss-Prot ver 14) shows it to be present only in cytochrome P-450 proteins.

(c) Isolation and characterization of *gfp* genomic clones

The *gfp1* cDNA was also used to isolate genomic clones prior to the availability of the *gfp10* cDNA. An *A. victoria* genomic library was constructed in λ2001 (Kam et al., 1984) essentially as described (Maniatis et al., 1982). Eight recombinant phages hybridizing to the *gfp1* cDNA were

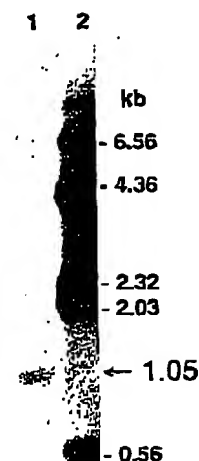


Fig. 3.

observed in GFP-derived peptides which showed a Tyr at position 100, a Met at position 141, but a Thr at position 103. Eight additional nt differences occur with the *gfp2* gene in the 3'-non-translated region of the *gfp10* cDNA (data not shown). It is not known whether the *gfp10* cDNA represents an allele of *gfp2* or another *gfp* gene.

These results will enable us to construct an expression vector for the preparation of non-fluorescent apoGFP. Since no information is yet available regarding the biosynthesis of the chromophore, a recombinant form of this protein will be a valuable reagent with which to examine the biochemistry of chromophore formation in this unique class of proteins and the mechanism of energy transfer between aequorin and GFP.

ACKNOWLEDGEMENTS

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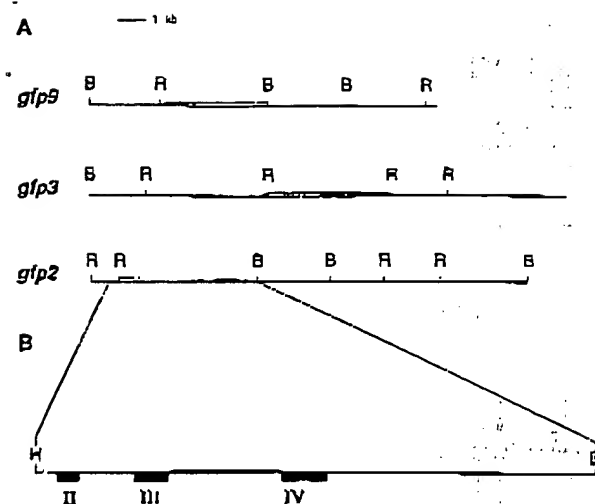


Fig. 4. Restriction enzyme maps of three *Aequorea gfp* genes. (A) The maps of three representative genomic clones are compared. The double lines represent those DNA fragments which hybridize to *gfp1* cDNA. Southern-blot analysis indicated three other genomic clones, λ GFP1, 4 and 8 (not shown) lack the 3' end of the gene. (B) The exon/intron arrangement of the gene encoded by λ GFP2 was determined by comparing the nt sequences of the 5-kb *EcoRI-BamHI* and the overlapping 1.8-kb *HindIII* fragments of λ GFP2 and the *EcoRI* insert of λ GFP10 cDNA. The exons are represented by the blackened boxes, I, II and III. The GenBank accession No. for the *gfp2* sequence is M62653.

purified from the genomic DNA library. Based on restriction enzyme and Southern-blot analyses, they represent six different isolates having at least three different restriction maps (Fig. 4). When DNA fragments from the 5'- and 3'-ends of the *gfp1* cDNA were used as hybridization probes, all of the genomic clones were found likely to contain the 5'-end of the gene, but only *gfp2*, 3, and 9 also contained the 3' end. The three types of genomic clones are consistent with the presence of multiple GFP isoforms isolated from *A. victoria* (A. Roth, M. Cutler and W.W.W., unpublished). Since the *A. victoria* genomic DNA used for the genomic library was isolated from a large number of

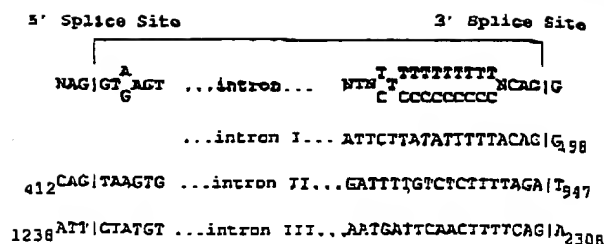


Fig. 5. Alignment of the nt sequences in *gfp2* at the splice junctions. The intron sequences were identified by comparing the nt sequences of *gfp2* and the *gfp10* cDNA (Fig. 2). The consensus sequence is taken from Senupathy et al. (1990).

jellyfish (collected at Friday Harbor, Washington), the three *gfp* genes are representative of the *Aequorea* population as opposed to individual jellyfish.

The *EcoRI-BamHI* and an overlapping *HindIII* fragments in the genomic clone λ GFP2 (Fig. 4) were sequenced and compared to that of the *gfp10* cDNA to examine the structure of the gene. The *gfp* gene encoded by λ GFP2 contains at least three exons spread over 2.6 kb of DNA (Fig. 4). These exons, designated II, III, and IV, encode 69, 98, and 71 aa, respectively. Presumably, a fourth exon is located upstream from the genome since the 15 nt at the 5' end of the *gfp10* cDNA sequence cannot be aligned to the 5' region of the DNA sequence derived from the *gfp2* gene. The positions of the introns with respect to the cDNA sequence are indicated (Fig. 2). The aa residues involved in the chromophore are encoded at the 3' end of exon II. The nt sequences of the *gfp* mRNA splice junctions agree reasonably well with consensus sequences (Fig. 5).

The *gfp10* cDNA is not encoded by the *gfp2* gene since there are several nt differences between their sequences. The nt differences within the protein-coding regions are summarized in Table 1A. Four of the 12 single nt differences result in conservative aa replacements at positions 100, 108, 141 and 219 (Table 1B). The aa residues encoded at these four positions are consistent with the aa sequences

TABLE 1

Sequence differences in the coding regions of the *gfp* clones

A	Nucleotide differences with respect to the <i>gfp2</i> gene ^a	B	Amino acid differences ^b		
			aa position	<i>gfp2</i> gene	<i>gfp10</i> cDNA <i>gfp1</i> cDNA
	<i>gfp10</i> cDNA	12 (8 silent)			
	<i>gfp1</i> cDNA	2 (2 silent)			
			100	Tyr	Phe
			108	Ser	Thr
			141	Met	Leu
			219	Ile	Val

^a Total number observed upon comparison of the nt sequences of the ORFs in the *gfp* cDNAs with the homologous sequences in the *gfp2* gene.

^b Observed upon comparison of the translations of the ORFs of both cDNAs and the exons of the *gfp2* gene. The aa numbering is the same as that used in Fig. 2.

United States Patent [19]

Halbritter

US005246631A

[11] Patent Number: 5,246,631

[45] Date of Patent: Sep. 21, 1993

[34] SELF-ILLUMINATED BUBBLES

[76] Inventor: Martin J. Halbritter, Holy Bakery
Rd., Kapaa, HI. 96755

[21] Appl. No.: 704,776

[22] Filed: May 23, 1991

[31] Int. Cl.⁵ C09K 3/00; A63H 33/28

[52] U.S. Cl. 252/700; 446/15

[58] Field of Search 252/582, 700; 446/15,
446/16, 17, 18, 19; 20, 21

[56] References Cited

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4,462,931	7/1984	Cohen et al.	252/700
4,717,511	1/1988	Koroscil	252/700

Primary Examiner—Philip Tucker

Attorney, Agent, or Firm—Pons, Smith, Lande & Rose

[37] ABSTRACT

Self-illuminated bubbles are disclosed wherein the bubbles are formed utilizing a bubble blowing solution which includes a sufficient amount of a surface active agent to form the bubbles and a sufficient amount of a chemiluminescent agent to provide self-illumination of the bubbles. The bubble blowing solution includes a surface active agent to provide formation of the bubble and a chemiluminescent agent such as CYALUME®. A kit for preparing the self-illuminated bubble blowing solution is also disclosed.

7 Claims, No Drawings

5,246,631

1

SELF-ILLUMINATED BUBBLES

BACKGROUND OF THE INVENTION

1. Field of the Invention:

The present invention relates generally to bubbles which are formed for recreational and demonstrative purposes. More particularly, the present invention relates to providing solutions which can be utilized to form self-illuminated bubbles.

2. Description of Related Art:

The formation of bubbles is a well recognized and widely practiced pastime. In its simplest form, bubble blowing involves dipping a ring-shaped article into a liquid soap solution followed by blowing into the ring to form one or more bubbles. Alternatively, the ring may be moved in order to force air through the ring and form the bubbles.

A wide variety of commercial bubble blowing kits are available for use by individuals. The kits typically include a bubble blowing solution packaged in a suitable container and some type of ring-shaped device for use in forming the bubbles. As an alternative to commercially available bubble blowing kits, individuals commonly practice the art of blowing bubbles utilizing liquid detergents, such as liquid dishwashing, and home-made ring-shaped devices which can be made from a variety of items.

The blowing of bubbles outdoors for recreational and entertainment purposes is limited, for the most part, to daylight hours. Although bubbles can be blown in the dark, the recreational and entertainment value drops substantially since it is difficult to see them. Likewise, bubble blowing indoors is limited to rooms where there is sufficient light to allow visual observation of the bubbles. Accordingly, it would be desirable to provide self-illuminated bubbles which can be viewed under conditions of reduced lighting. The self-illumination of bubbles would also increase the entertainment value and enjoyment of the bubbles under conditions when the bubbles would otherwise be difficult to see. In addition, the self-illumination of bubbles would provide a source of light and illumination in an attractive manner.

SUMMARY OF THE INVENTION

In accordance with the present invention it was discovered that self-illuminated bubbles can be formed which substantially increase the visibility of bubbles when blown in the dark. The self-illuminated bubbles in accordance with the present invention not only are more visible in the dark, but also provide an attractive and appealing source of light. The self-illuminated bubble emits light continuously as the bubble is formed and during the bubbles life as it floats through the air and eventually collapses.

The present invention is based upon the discovery that chemiluminescent agents can be added to conventional bubble blowing solutions in amounts which are sufficient to provide self-illumination of the bubbles without deleteriously affecting the capability of the solution to form suitable bubbles. A bubble blowing solution in accordance with the present invention includes a sufficient amount of a surface-active agent to provide formation of the bubbles when a gas is introduced into contact with the solution and a sufficient amount of a chemiluminescent agent to provide self-illumination of the bubbles. As a feature of the present invention, the chemiluminescent agent is an oxalate

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diester which reacts with a peroxide and a fluorescer to provide the self-illumination of the bubble. The chemiluminescent agent, such as CYALUME® available from American Cyanamide, can be added to the bubble blowing solution in amounts up to about 20 volume percent.

As a feature of the present invention, it was discovered that bubble hardeners such as glycerin may also be added to the bubble blowing solution in amounts up to about one volume percent to enhance bubble formation. It was discovered that addition of the bubble hardeners does not adversely affect the bubble formation or self-illumination properties of the bubble blowing solution in accordance with the present invention.

The present invention provides a simple and effective method for illuminating bubbles for use in a wide variety of educational, entertainment and recreational settings. The self-illuminated bubbles in accordance with the present invention provide an especially attractive bubble which provides a unique visual glow in the dark. The above-discussed and many other features and attendant advantages of the present invention will become better understood by reference to the following detailed description.

DETAILED DESCRIPTION OF THE INVENTION

The present invention involves self-illuminated bubbles and the solutions used in their preparation. The bubble blowing solutions in accordance with the present invention have wide application for use in a variety of bubble blowing devices. The self-illuminated bubble blowing solution of the present invention may be used in simple ring-shaped bubble blowing devices intended for personalized use. The self-illuminated bubble blowing solution also is suitable for the blowing of bubbles on a larger scale by mechanized equipment.

The present invention is based upon the addition of a chemiluminescent agent to an otherwise conventional bubble blowing solution. The chemiluminescent agent can be added to a wide variety of bubble blowing solutions provided that there is no deleterious reaction between the chemiluminescent agent and the bubble blowing solution. Suitable bubble blowing solutions may include anionic, cationic, non-ionic and ampholytic surfactants. Many of the conventional solutions utilized in bubble blowing kits or in conjunction with bubble blowing machinery are suitable provided that a sufficient amount of a surface active agent is included to provide formation of bubbles when a gas is introduced into contact with the solution.

Solutions containing conventional surfactants, such as sodium laureth sulfate or ammonium laureth sulfate may be utilized. Particularly preferred bubble blowing solutions are Liquid s for dishes. The liquid may be used full-strength or may be diluted with water to provide the desired consistency of bubbles depending upon the types of bubbles desired. The amount of water, if any, used to dilute the liquid is not particularly critical and can be varied as is well-known in the art to achieve desired bubble consistency.

The chemiluminescent agent which is added to the bubble blowing solution in accordance with the present invention may be selected from a wide variety of light producing materials. However, the preferred chemiluminescent agent includes an oxalate diester which reacts with a peroxide and a fluorescer to provide the emission

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of light. This particular light emitting compound and chemiluminescent reaction is well-known and forms the basis for a variety of chemiluminescent compositions.

The composition and chemistry of exemplary chemiluminescent agents is described in detail in U.S. Pat. Nos. 3,749,679; 3,775,336; 3,888,786; 3,911,038; 4,017,415; 4,076,645; 4,313,843; 4,379,320; 4,768,608; and 4,717,511. These patents are all assigned to American Cyanamide Company (Stanford, Connecticut) and are hereby incorporated by reference. A particularly preferred chemiluminescent agent for use in accordance with the present invention is Cyalume® which is a commercially available product which is marketed by American Cyanamide and is covered by the above-identified patents. CYALUME® is available as a kit which can be mixed together following the manufacturers instructions to form the solution which is chemiluminescent.

Bubble hardeners such as glycerin may also be added to the bubble blowing solution in accordance with the present invention. Up to one percent by volume bubble hardener may be added to the solution with approximately 0.2 percent being preferred.

As is well-known, oxalate diesters do not begin to emit light until they are mixed with peroxide and a fluorescer. Once the reaction begins, the period over which light will be emitted can range from a few minutes to a few hours. Accordingly, it is preferred, in accordance with the present invention, that the combination of oxalate diester with a peroxide and a fluorescer be delayed until just prior to use in the bubble blowing solution. Preferably, a bubble blowing kit is provided wherein the bubble blowing solution comprising the surface active agent is packaged separately from the chemiluminescent agent. The kit includes instructions for mixing the chemiluminescent materials together to form the chemiluminescent agent. The instructions further describe the process for mixing the chemiluminescent agent with the bubble blowing solution in order to make the solution light emitting. Alternatively, the bubble blowing solution may contain the oxalate diester with the peroxide and fluorescer being packaged separately.

The amount of chemiluminescent agent which can be added to the bubble blowing solution may be varied in order to provide the desired degree of illumination. Preferably, the amount of chemiluminescent agent will not exceed about 20 volume percent. Larger amounts of chemiluminescent agent do not substantially increase

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the amount of light emitted and also tend to reduce bubble quality.

An example of practice of the present invention involves using a liquid dish such as LEMON JOY® available from Procter & Gamble Company (Cincinnati, Ohio). This particular dishwashing liquid is covered by U.S. Pat. Nos. 4,133,779 and 4,316,824, the contents of which are hereby incorporated by reference. Although the LEMON JOY® may be diluted with varying amounts of water, it is preferred that the dishwashing liquid be used at full strength. Approximately 9 milliliters of CYALUME® solution made in accordance with the manufacturers instructions are added to approximately 120 milliliters of the dishwashing liquid. Although this particular mixture may be used to produce adequate self-illuminated bubbles, it is preferred that 3 to 4 drops of glycerin be added to the solution as a bubble hardener. The solution is then ready for use to form self-illuminated bubbles.

Having thus described exemplary embodiments of the present invention, it should be noted by those skilled in the art that the within disclosures are exemplary only and that various other alternatives, adaptations and modifications may be made within the scope of the present invention. Accordingly, the present invention is not limited to the specific embodiments as illustrated herein, but is only limited by the following claims.

What is claimed is:

1. A self-illuminated bubble comprising a solution containing a sufficient amount of a surface active agent to provide formation of said bubble, and a sufficient amount of a chemiluminescent agent to provide self-illumination of said bubble.

2. A self-illuminated bubble according to claim 1 wherein said solution is an aqueous solution.

3. A self-illuminated bubble according to claim 2 wherein said surface active agent is selected from the group of surface active agents consisting of anionic, cationic, non-ionic and amphoteric surfactants.

4. A self-illuminated bubble according to claim 1 wherein said chemiluminescent agent comprises an oxalate diester which reacts with a peroxide and a fluorescer to provide said self-illumination of said bubble.

5. A self-illuminated bubble according to claim 4 wherein said chemiluminescent agent comprises an oxalate diester, said peroxide and said fluorescer.

6. A self-illuminated bubble according to claim 2 wherein said surface active agent is a liquid soap.

7. A self-illuminated bubble according to claim 6 wherein said aqueous solution further comprises less than about 1 volume percent glycerin.

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